

# Introduction to computers and computing environments

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Math  
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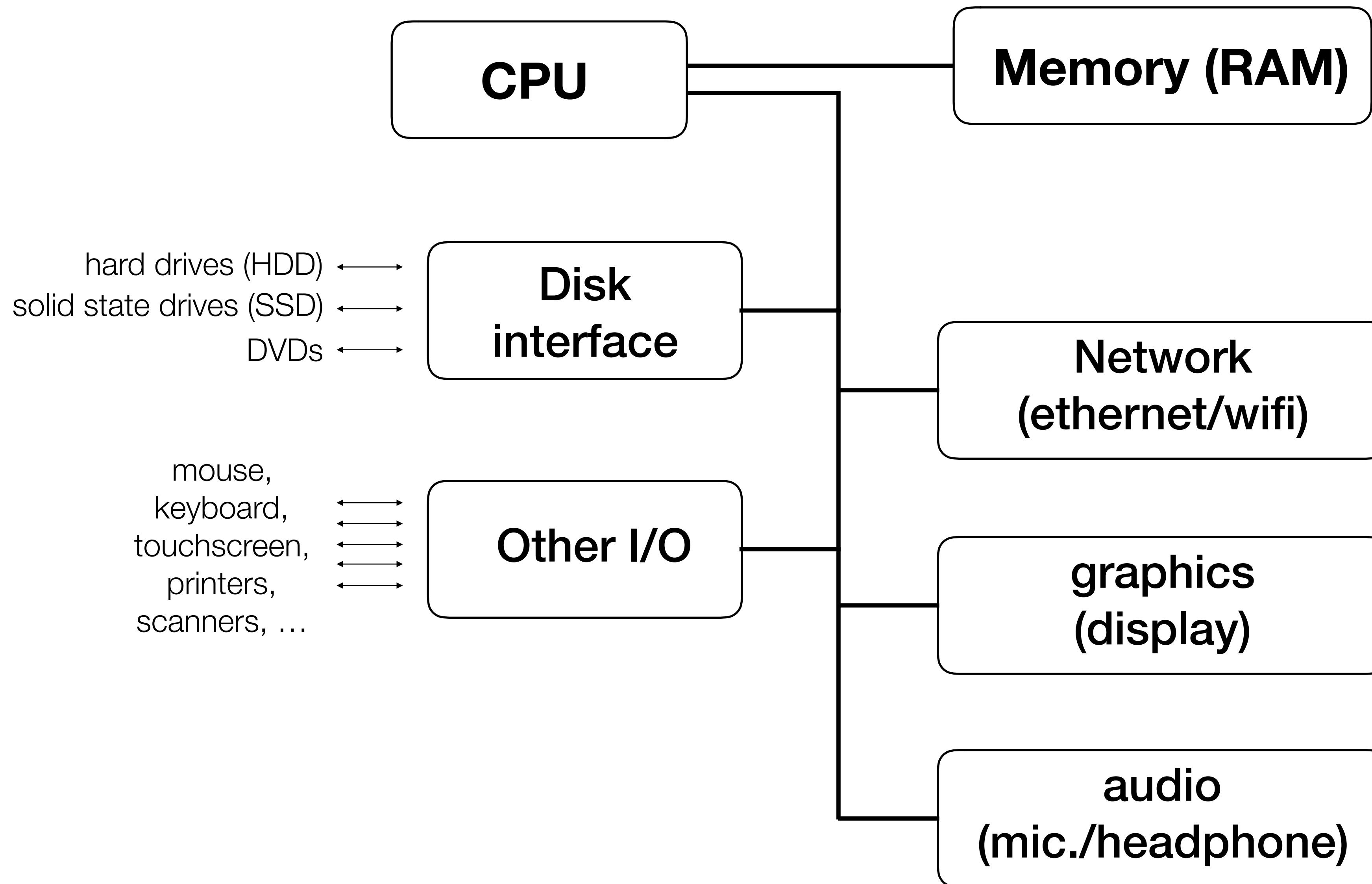
The Arthropod-Borne and  
Infectious Diseases Laboratory  
3185 Rampart Road Building T

AIDL  
Colorado State  
University

Director, CSU  
NGS facility



# Basic computer architecture



# Memory (RAM) vs. storage (drives)

	<b>Hard drives (HDD)</b>	<b>Solid state drives (SSD)</b>	<b>Main memory (RAM)</b>
Speed	Slow	Much faster	Fastest
Cost per Gb	\$0.025	\$0.20	\$4
Volatile	No	No	Yes
Uses	Long term storage of large datasets	Laptops, long term storage, OS boot drives	
Other comments	Uses more power	Largest SSD ~4 Tb	Some bioinformatic tasks, like genome assembly, require lots of RAM



# CPUs vs. cores vs. threads

Most modern computers have more than one CPU

Typically, these are referred to as “cores”.



To make it more confusing, Intel cores are “hyper-threaded”, which means that the operating systems sees these 4 cores as 8 CPUs.

```
[MDSTENGL-M01:Desktop _mdstengl$ sysctl hw.physicalcpu hw.logicalcpu
hw.physicalcpu: 4
hw.logicalcpu: 8
```

# Having 8 CPUs on your laptops means you can run programs in parallel

map 1M 50 nt reads to the human genome with 1 CPU: 1m 16 seconds

```
mdstengl@cctsi-104:~/analyses/test_human_mapping$ time ./run_bowtie ERR3252925_1_1M.fastq GCA_000001405.15_GRCh38_no_alt_analysis_set.fna.bowtie_index
bowtie2 -x GCA_000001405.15_GRCh38_no_alt_analysis_set.fna.bowtie_index -q -U ERR3252925_1_1M.fastq --local --score-min C,1
00,1 --no-unal --threads 1 -S ERR3252925_1_1M.fastq.GCA_000001405.15_GRCh38_no_alt_analysis_set.fna.bowtie_index.sam

real    1m16.427s
user    1m13.836s
sys     0m12.844s
```

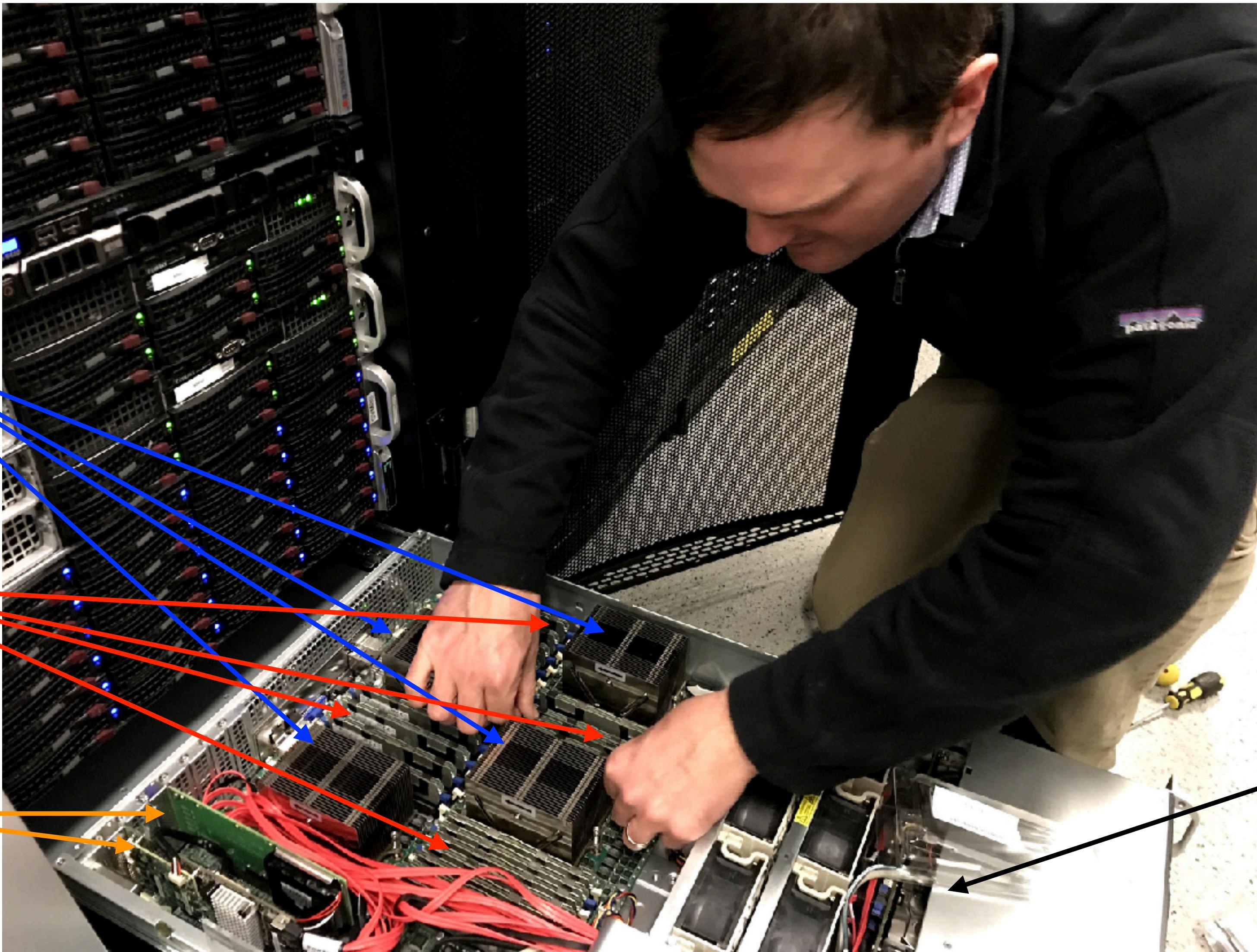
map 1M 50 nt reads to the human genome with 24 CPU: 9.6 seconds

```
mdstengl@cctsi-104:~/analyses/test_human_mapping$ time ./run_bowtie_multiple_threads ERR3252925_1_1M.fastq GCA_000001405.15
_GRCh38_no_alt_analysis_set.fna.bowtie_index
bowtie2 -x GCA_000001405.15_GRCh38_no_alt_analysis_set.fna.bowtie_index -q -U ERR3252925_1_1M.fastq --local --score-min C,1
00,1 --no-unal --threads 24 -S ERR3252925_1_1M.fastq.GCA_000001405.15_GRCh38_no_alt_analysis_set.fna.bowtie_index.sam

real    0m9.641s
user    1m38.696s
sys     0m33.124s
```

Taking advantage of multiple CPUs is called multi-processing (or multi-threading)  
When possible, you should take advantage of available cores.

# A bioinformatics server



CPUs

( $4 \times 16$  cores =  
64)

RAM

( $16 \times 32\text{Gb} =$   
512 Gb)

ethernet

(main user  
interface)

HDDs ( $6 \times 8$  Tb)

& SSDs ( $2 \times 128\text{Gb}$ )

RAID configured

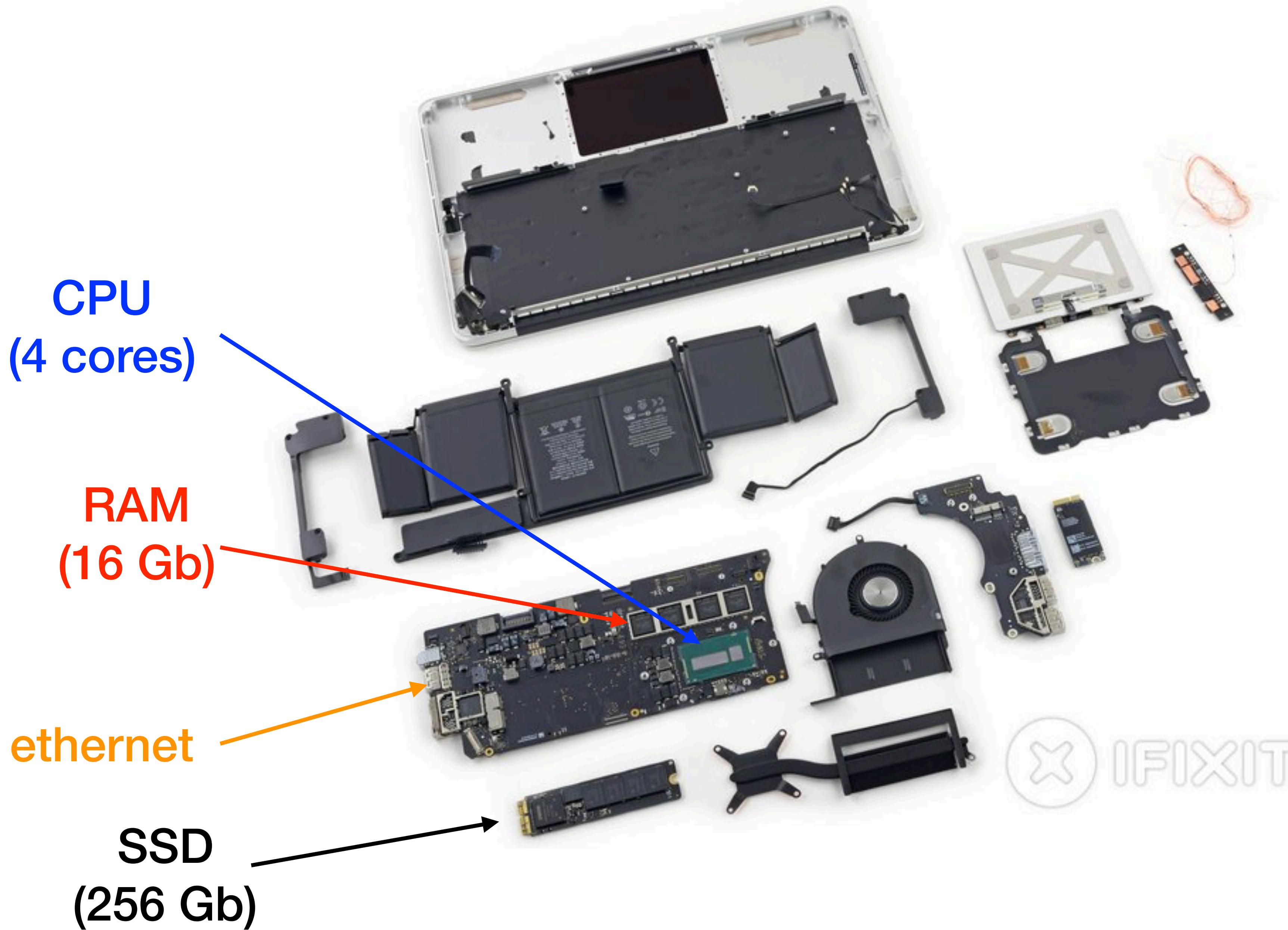
# How do you check resource usage and availability?

## **On your workshop laptops:**

- How much memory is available? How much is used?
- How much storage is available? What type is it? How much is used?
- How many cores and threads are there? What fraction of these are being used?



# The inside of your workshop laptops



# Checking resource usage and availability through the command line

command	better way to run it
df	reports disk usage and storage
top	reports CPU & memory usage

## Use these commands to answer the same questions:

- How much storage is available? How much is used?
- How much memory is available? How much is used?
- How many cores and threads are there? What fraction of these are being used?

# Local vs. remote computing

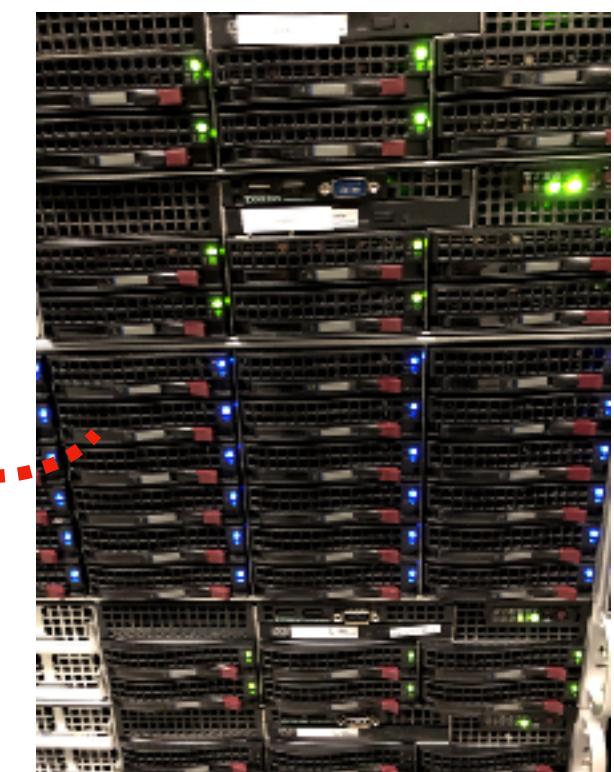
Local: you are using the resources  
on your computer



Remote: you are using the resources of a  
distant computer, but probably still  
connected through your own computer



a server  
somewhere



# Let's connect to a remote server!

**Connect to a bioinformatics server at CSU: Open the terminal app and run:**

```
ssh gdw@cctsi-104.cvmbs.colostate.edu
```

**Use the df and top commands to answer the same questions about this server**

- How much storage is available? How much is used? (hint: in /home)
- How much memory is available? How much is used?
- How many cores are there? What fraction of these are being used?

# Cloud computing is an increasingly popular form of remote computing

## Advantages:

- Scalable and flexible
- Don't have to buy or maintain servers
- Can take advantage of pre-existing images and containers

## Disadvantages:

- Can be expensive
- Have to pay to store and transfer data
- Can be slow to transfer data

Real physical computing resources are available as virtual computers through the internet

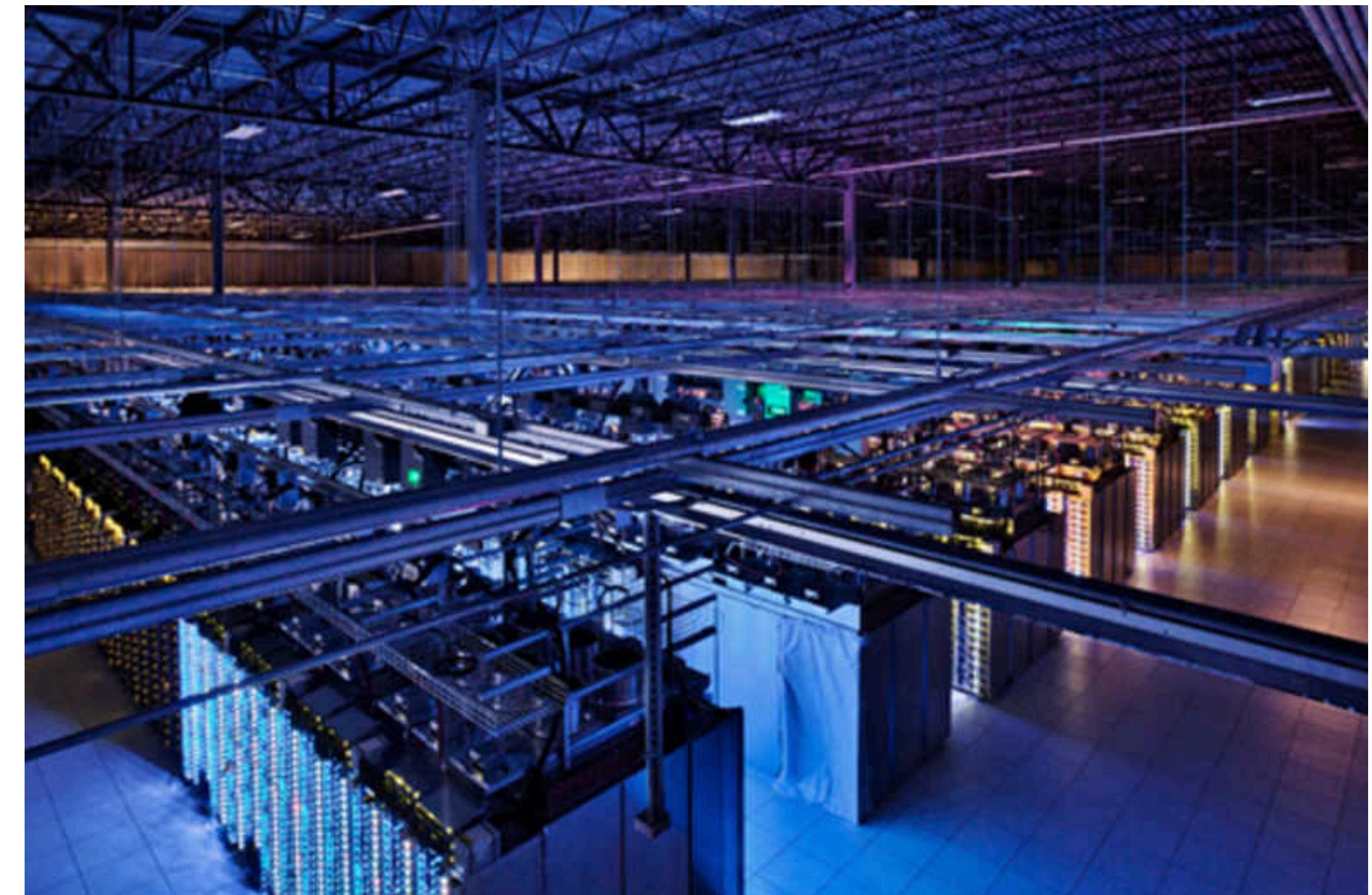


image: gigabitmagazine.com

# Cost of cloud computing depends on how many resources you need

	vCPU	ECU	Memory (GiB)	Instance Storage (GB)	Linux/UNIX Usage
t2.large	2	Variable	8 GiB	EBS Only	\$0.1104 per Hour
t2.xlarge	4	Variable	16 GiB	EBS Only	\$0.2208 per Hour
r5.12xlarge	48	173	384 GiB	EBS Only	\$3.36 per Hour
r5.24xlarge	96	347	768 GiB	EBS Only	\$6.72 per Hour

Similar to the workshop laptops

Similar to the servers my lab uses

Plus ~\$0.05-\$0.15 per Gb-month for (short term) storage: \$50-\$150 per month per Tb

Plus ~\$0.09 per Gb to transfer from Amazon to anywhere else: \$90 per Tb

# An Amazon Web Services (AWS) linux environment

The screenshot shows the AWS EC2 console interface. On the left, the navigation menu includes options like EC2 Dashboard, Instances (selected), Launch Templates, Spot Requests, Reserved Instances, Dedicated Hosts, Scheduled Instances, Capacity Reservations, AMIs, and more. In the center, a terminal window displays a root shell session on an Ubuntu 18.04.2 LTS instance (ami-094d903b47d77bec). The session shows system information, package updates, and a login history. On the right, a sidebar shows the instance's details, including its private DNS name (ip-172-31-49-140.us-west-2.compute.internal), private IP (172.31.49.140), VPC ID (vpc-bfb11c7), and subnet ID (subnet-c7d3f28c).

ubuntu@ip-172-31-49-140:~\$

Pv6 IPs	Key Name	Monitoring	Launch Time
	mds_linux	disabled	May 29, 2019 at 10:37:
	mds_linux	disabled	May 29, 2019 at 10:42:

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